



## SEQUENCE LISTING

<110> Mahajan, Muktar A.  
Samuels, Herbert H.

<120> NIF-1 IS A NOVEL CO-TRANSDUCER THAT INTERACTS WITH AND  
REGULATES THE ACTIVITY OF THE NUCLEAR HORMONE RECEPTOR  
CO-ACTIVATOR, NRC

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<151> 2002-08-23

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<170> PatentIn Ver. 2.1

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			275		280						285				
Ser	Ala	Gln	Arg	Glu	Leu	Gly	Ala	Thr	Ala	Leu	Gln	Val	Ala	Val	Val
			290		295						300				
Lys	Ser	Glu	Asp	Val	Glu	Ala	Glu	Leu	Thr	Ser	Thr	Ala	Arg	Gln	Pro
			305		310				315						320
Ser	Ser	Glu	Asp	Thr	Thr	Pro	Arg	Val	Val	Thr	Leu	His	Val	Ala	Glu
			325			330									335
Ser	Gly	Ser	Ser	Val	Ala	Ala	Glu	Ser	Gln	Leu	Gly	Pro	Ser	Asp	Leu
			340			345									350
Gln	Gln	Ile	Ala	Leu	Pro	Pro	Gly	Pro	Phe	Ser	Gly	Ala	Ser	Tyr	Ser
			355		360						365				
Val	Ile	Thr	Ala	Pro	Pro	Val	Glu	Gly	Arg	Ala	Ser	Ala	Ser	Gly	Pro
			370		375						380				
Pro	Tyr	Arg	Glu	Glu	Pro	Pro	Gly	Glu	Ala	Ala	Gln	Ala	Val	Val	Val
			385		390				395						400
Asn	Asp	Thr	Leu	Lys	Glu	Ala	Gly	Thr	His	Tyr	Ile	Met	Ala	Ala	Asp
			405				410								415

Gly Thr Gln Leu His His Ile Glu Leu Thr Ala Asp Gly Ser Ile Ser  
                  420                        425                        430  
  
 Phe Pro Ser Pro Asp Thr Leu Ala Pro Gly Thr Lys Trp Pro Leu Leu  
                  435                        440                        445  
  
 Gln Cys Gly Gly Pro Pro Arg Asp Gly Pro Glu Val Leu Ser Pro Thr  
                  450                        455                        460  
  
 Lys Thr His His Thr Gly Gly Ser Gln Gly Ser Ser Thr Pro Pro Pro  
                  465                        470                        475                        480  
  
 Ala Thr Ser His Ala Leu Gly Leu Leu Val Pro His Ser Pro Pro Ser  
                  485                        490                        495  
  
 Ala Ala Ala Ser Ser Thr Lys Lys Phe Ser Cys Lys Val Cys Ser Glu  
                  500                        505                        510  
  
 Ala Phe Pro Ser Arg Ala Glu Met Glu Ser His Lys Arg Ala His Ala  
                  515                        520                        525  
  
 Gly Pro Ala Ala Phe Lys Cys Pro Asp Cys Pro Phe Ser Ala Arg Gln  
                  530                        535                        540  
  
 Trp Pro Glu Val Arg Ala His Met Ala Gln His Ser Ser Leu Arg Pro  
                  545                        550                        555                        560  
  
 His Gln Cys Asn Gln Cys Ser Phe Ala Ser Lys Asn Lys Lys Asp Leu  
                  565                        570                        575  
  
 Arg Arg His Met Leu Thr His Thr Asn Glu Lys Pro Phe Ser Cys His  
                  580                        585                        590  
  
 Val Cys Gly Gln Arg Phe Asn Arg Asn Gly His Leu Lys Phe His Ile  
                  595                        600                        605  
  
 Gln Arg Leu His Ser Ile Asp Gly Arg Lys Thr Gly Thr Ser Thr Ala  
                  610                        615                        620  
  
 Arg Ala Pro Ala Gln Thr Ile Ile Leu Asn Ser Glu Glu Glu Thr Leu  
                  625                        630                        635                        640  
  
 Ala Thr Leu His Thr Ala Phe Gln Ser Asn His Gly Thr Leu Gly Thr  
                  645                        650                        655  
  
 Glu Arg Leu Gln Gln Ala Leu Ser Gln Glu His Ile Ile Val Ala Gln  
                  660                        665                        670

Glu Gln Thr Val Ala Asn Gln Glu Glu Ala Thr Tyr Ile Gln Glu Ile  
675 680 685

Thr Ala Asp Gly Gln Thr Val Gln His Leu Val Thr Ser Asp Asn Gln  
690 695 700

Val Gln Tyr Ile Ile Ser Gln Asp Gly Val Gln His Leu Leu Pro Gln  
705 710 715 720

Glu Tyr Val Val Val Pro Asp Gly His His Ile Gln Val Gln Glu Gly  
725 730 735

Gln Ile Thr His Ile Gln Tyr Glu Gln Gly Thr Pro Phe Leu Gln Glu  
740 745 750

Ser Gln Ile Gln Tyr Val Pro Val Ser Pro Ser Gln Gln Leu Val Thr  
755 760 765

Gln Ala Gln Leu Glu Ala Ala Ala His Ser Ala Val Thr Val Ala Asp  
770 775 780

Ala Ala Met Ala Gln Ala Gln Gly Leu Phe Gly Thr Glu Glu Ala Val  
785 790 795 800

Pro Glu His Ile Gln Gln Leu Gln His Gln Gly Ile Glu Tyr Asp Val  
805 810 815

Ile Thr Leu Ser Asp Asp  
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Ala Val Asn Ala Ala

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<211> 122

<212> PRT

<213> Human

<400> 12

Cys Asp Lys Cys Gly Lys Ser Phe Lys Lys Arg Tyr Thr Phe Lys Met  
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His Leu Leu Thr His Cys Glu Phe Val Cys Glu Asp Lys Lys Ala Leu  
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Leu Asn His Gln Leu Ser His Ala Thr Gln Thr Ala Leu Asp Leu Leu  
35 40 45

Leu Asn Met Ser Ala Gln Arg Glu Leu Cys Lys Ile Cys Ala Glu Ala  
50 55 60

Phe Pro Gly Arg Ala Glu Met Glu Ser His Lys Arg Ala His Cys His  
65 70 75 80

Leu Cys Gly Gln Arg Phe Asn Arg Asn Gly His Leu Lys Phe His Ile  
85 90 95

Gln Arg Leu His Leu Asn Ser Asp Asp Glu Thr Leu Ala Thr Leu His  
100 105 110

Thr Ala Leu Gln Ser Ser His Gly Val Leu  
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Arg Asn Gly Glu Ser Gln Glu Ser His Gln Ile Met Glu Asp Gln Gly  
20 25 30  
  
Gln Ala

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Val Ser Ser Val Ile Glu Glu Glu Phe Asn Thr  
1 5 10